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SEQUENCE LISTING



<110> TORDO, NOEL  
PERRIN, PIERRE  
JACOB, YVES  
BAHLOUL, CHOKRI

<120> CHIMERIC LYSSAVIRUS NUCLEIC ACIDS AND POLYPEPTIDES

<130> 03495-0188-00000

<140> 09/549,519  
<141> 2000-04-14

<150> 60/129,501  
<151> 1999-04-15

<160> 40

<170> PatentIn Ver. 2.1

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<223> Description of Artificial Sequence: Primer

<400> 1

ttcttagagcc accatgggttc ctcaggctct cctg

34

<210> 2  
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<223> Description of Artificial Sequence: Primer

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attgatcaac tgaccgggag ggc

23

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<211> 98  
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<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic  
oligonucleotide PVp1

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aattcttagag cccgccaccat ggttcctcag gctctcctgt ttgtacccct tctggttttt 60  
ccattgtgtt ttgggaagaa ttcccccccc ggtcagtt 98

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oligonucleotide PVp2

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gatcaactga ccgggggggg aattttccc aaaacacaat ggaaaaacca gaaggggtac 60  
aaacaggaga gcctgaggaa ccatggtggc ggctctag 98

<210> 5  
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oligonucleotide EBL1p1

<400> 5  
aatttcccaa tctacaccat cccggataaa atcggaccgt ggtcacctat tccg 54

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oligonucleotide EBL1p2

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aattcggaat aggtgaccac ggtccgattt tatccggat ggttagatt ggga 54

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gag 63

<210> 8  
<211> 27  
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&lt;220&gt;

&lt;223&gt; Description of Artificial Sequence: Primer

&lt;400&gt; 8

ggaattcgag caccattctg' gagcttc

27

&lt;210&gt; 9

&lt;211&gt; 11

&lt;212&gt; PRT

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; Description of Artificial Sequence: Synthetic peptide

&lt;400&gt; 9

Asp Asn Pro Ala Ser Thr Thr Asn Lys Asp Lys

1

5

10

&lt;210&gt; 10

&lt;211&gt; 9

&lt;212&gt; PRT

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; Description of Artificial Sequence: Synthetic peptide

&lt;400&gt; 10

Pro Gln Ala Ser Gly Val Tyr Met Gly

1

5

&lt;210&gt; 11

&lt;211&gt; 16

&lt;212&gt; PRT

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; Description of Artificial Sequence: Synthetic peptide

&lt;400&gt; 11

Glu Arg Pro Gln Ala Ser Gly Val Tyr Met Gly Asn Leu Thr Ala Gln

1

5

10

15

&lt;210&gt; 12

&lt;211&gt; 78

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; Description of Artificial Sequence: Primer

<400> 12  
aattcagata acccggcgta gaccactaac aaggataagc tgttcgagt gcctcaggcc 60  
tctgggtgt atatgggt 78

<210> 13  
<211> 78  
<212> DNA  
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<223> Description of Artificial Sequence: Primer

<400> 13  
aattacccat atacacacca gaggcctgag gcactgcgaa cagcttatcc ttgttagtgg 60  
tcgacgcccgg gttatctg 78

<210> 14  
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<223> Description of Artificial Sequence: Synthetic  
adaptor

<400> 14  
aatttggata acccggcgta gaccactaac aa 32

<210> 15  
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<212> DNA  
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<223> Description of Artificial Sequence: Synthetic  
adaptor

<400> 15  
aattcttatac cttgttagtg gtcgacgcccgg 32

<210> 16  
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<212> DNA  
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<223> Description of Artificial Sequence: Synthetic  
adaptor

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<210> 17

<211> 54  
 <212> DNA  
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<220>  
 <223> Description of Artificial Sequence: Synthetic adaptor

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 aattctggga agtaagatta cccatataca caccagaggc ctgaggtctc tcca

54

<210> 18  
 <211> 61  
 <212> PRT  
 <213> Artificial Sequence

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 <223> Description of Artificial Sequence: Chimeric polypeptide PV-PV

<400> 18  
 Val Leu Gly Leu Arg Leu Met Asp Gly Thr Trp Val Ser Met Gln Thr  
 1 5 10 15

Ser Asn Glu Thr Lys Trp Cys Pro Pro Gly Gln Leu Ile Asn Leu His  
 20 25 30

Asp Phe Arg Ser Asp Glu Ile Glu His Leu Val Val Glu Glu Leu Val  
 35 40 45

Lys Lys Arg Glu Glu Cys Leu Asp Ala Leu Glu Ser Ile  
 50 55 60

<210> 19  
 <211> 39  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> Description of Artificial Sequence: Chimeric polypeptide -PVIII

<400> 19  
 Ser Pro Pro Gly Gln Leu Ile Asn Leu His Asp Phe Arg Ser Asp Glu  
 1 5 10 15

Ile Glu His Leu Val Val Glu Leu Val Lys Lys Arg Glu Glu Cys  
 20 25 30

Leu Asp Ala Leu Glu Ser Ile  
 35

<210> 20  
 <211> 64  
 <212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chimeric polypeptide 'EBL1-PV'

<400> 20

Val	Pro	Gly	Met	Arg	Leu	Met	Asp	Gly	Ser	Trp	Val	Ser	Leu	Gln	Lys
1					5					10				15	

Thr	Glu	Ala	Pro	Glu	Trp	Cys	Ser	Asn	Ser	Pro	Pro	Gly	Gln	Leu	Ile
					20			25					30		

Asn	Leu	His	Asp	Phe	Arg	Ser	Asp	Glu	Ile	Glu	His	Leu	Val	Val	Glu
					35			40					45		

Glu	Leu	Val	Lys	Lys	Arg	Glu	Glu	Cys	Leu	Asp	Ala	Leu	Glu	Ser	Ile
					50			55				60			

<210> 21

<211> 61

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chimeric polypeptide Mok-PV

<400> 21

Arg	Pro	Gly	Ile	Arg	Leu	Phe	Asp	Gly	Thr	Trp	Val	Ser	Phe	Thr	Lys
1					5				10				15		

Pro	Asp	Val	His	Val	Trp	Cys	Thr	Pro	Asn	Gln	Leu	Ile	Asn	Leu	His
				20				25				30			

Asp	Phe	Arg	Ser	Asp	Glu	Ile	Glu	His	Leu	Val	Val	Glu	Glu	Leu	Val
					35			40				45			

Lys	Lys	Arg	Glu	Glu	Cys	Leu	Asp	Ala	Leu	Glu	Ser	Ile
					50			55				60

<210> 22

<211> 61

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chimeric polypeptide PV-Mok

<400> 22

Val	Leu	Gly	Leu	Arg	Leu	Met	Asp	Gly	Thr	Trp	Val	Ser	Met	Gln	Thr
1					5				10				15		

Ser	Asn	Glu	Thr	Lys	Trp	Cys	Pro	Pro	Gly	Gln	Leu	Ile	Asn	Ile	His
				20				25				30			

Asn Asp Arg Leu Asp Glu Ile Glu His Leu Ile Val Glu Asp Ile Ile  
 35 40 45

Lys Lys Arg Glu Glu Cys Leu Asp Thr Leu Glu Thr Ile  
 50 55 60

<210> 23

<211> 61

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Chimeric  
 polypeptide Mok-Sad

<400> 23

Lys Pro Gly Ile Arg Leu Phe Asp Gly Thr Trp Val Ser Phe Thr Lys  
 1 5 10 15

Pro Asp Glu Thr Lys Trp Cys Pro Pro Asp Lys Leu Val Asn Leu His  
 20 25 30

Asp Phe Arg Ser Asp Glu Ile Glu His Leu Val Val Glu Glu Leu Val  
 35 40 45

Arg Lys Arg Glu Glu Cys Leu Asp Ala Leu Glu Ser Ile  
 50 55 60

<210> 24

<211> 524

<212> PRT

<213> Lyssavirus sp.

<220>

<223> PV

<400> 24

Met Val Pro Gln Ala Leu Leu Phe Val Pro Leu Leu Val Phe Pro Leu  
 1 5 10 15

Cys Phe Gly Lys Phe Pro Ile Tyr Thr Ile Pro Asp Lys Leu Gly Pro  
 20 25 30

Trp Ser Pro Ile Asp Ile His His Leu Ser Cys Pro Asn Asn Leu Val  
 35 40 45

Val Glu Asp Glu Gly Cys Thr Asn Leu Ser Gly Phe Ser Tyr Met Glu  
 50 55 60

Leu Lys Val Gly Tyr Ile Ser Ala Ile Lys Met Asn Gly Phe Thr Cys  
 65 70 75 80

Thr Gly Val Val Thr Glu Ala Glu Thr Tyr Thr Asn Phe Val Gly Tyr  
 85 90 95

Val Thr Thr Phe Lys Arg Lys His Phe Arg Pro Thr Pro Asp Ala  
 100 105 110

Cys Arg Ala Ala Tyr Asn Trp Lys Met Ala Gly Asp Pro Arg Tyr Glu  
 115 120 125

Glu Ser Leu His Asn Pro Tyr Pro Asp Tyr His Trp Leu Arg Thr Val  
 130 135 140

Lys Thr Thr Lys Glu Ser Leu Val Ile Ile Ser Pro Ser Val Ala Asp  
 145 150 155 160

Leu Asp Pro Tyr Asp Arg Ser Leu His Ser Arg Val Phe Pro Gly Gly  
 165 170 175

Asn Cys Ser Gly Val Ala Val Ser Ser Thr Tyr Cys Ser Thr Asn His  
 180 185 190

Asp Tyr Thr Ile Trp Met Pro Glu Asn Pro Arg Leu Gly Met Ser Cys  
 195 200 205

Asp Ile Phe Thr Asn Ser Arg Gly Lys Arg Ala Ser Lys Gly Ser Glu  
 210 215 220

Thr Cys Gly Phe Val Asp Glu Arg Gly Leu Tyr Lys Ser Leu Lys Gly  
 225 230 235 240

Ala Cys Lys Leu Lys Leu Cys Gly Val Leu Gly Leu Arg Leu Met Asp  
 245 250 255

Gly Thr Trp Val Ala Met Gln Thr Ser Asn Glu Thr Lys Trp Cys Pro  
 260 265 270

Pro Gly Gln Leu Val Asn Leu His Asp Phe Arg Ser Asp Glu Ile Glu  
 275 280 285

His Leu Val Val Glu Glu Leu Val Lys Lys Arg Glu Glu Cys Leu Asp  
 290 295 300

Ala Leu Glu Ser Ile Met Thr Thr Lys Ser Val Ser Phe Arg Arg Leu  
 305 310 315 320

Ser His Leu Arg Lys Leu Val Pro Gly Phe Gly Lys Ala Tyr Thr Ile  
 325 330 335

Phe Asn Lys Thr Leu Met Glu Ala Asp Ala His Tyr Lys Ser Val Arg  
 340 345 350

Thr Trp Asn Glu Ile Ile Pro Ser Lys Gly Cys Leu Arg Val Gly Gly  
 355 360 365

Arg Cys His Pro His Val Asn Gly Val Phe Phe Asn Gly Ile Ile Leu  
 370 375 380

Gly Pro Asp Gly Asn Val Leu Ile Pro Glu Met Gln Ser Ser Leu Leu  
 385 390 395 400

Gln	Gln	His	Met	Glu	Leu	Leu	Val	Ser	Ser	Val	Ile	Pro	Leu	Met	His
														415	
405															
Pro	Leu	Ala	Asp	Pro	Ser	Thr	Val	Phe	Lys	Asn	Gly	Asp	Glu	Ala	Glu
														430	
420								425							
Asp	Phe	Val	Glu	Val	His	Leu	Pro	Asp	Val	His	Glu	Arg	Ile	Ser	Gly
													445		
435								440							
Val	Asp	Leu	Gly	Leu	Pro	Asn	Trp	Gly	Lys	Tyr	Val	Leu	Leu	Ser	Ala
													460		
450								455							
Gly	Ala	Leu	Thr	Ala	Leu	Met	Leu	Ile	Ile	Phe	Leu	Met	Thr	Cys	Trp
								470				475			480
465															
Arg	Arg	Val	Asn	Arg	Ser	Glu	Pro	Thr	Gln	His	Asn	Leu	Arg	Gly	Thr
													495		
485															
Gly	Arg	Glu	Val	Ser	Val	Thr	Pro	Gln	Ser	Gly	Lys	Ile	Ile	Ser	Ser
								500				505			510
Trp	Glu	Ser	Tyr	Lys	Ser	Gly	Gly	Glu	Thr	Gly	Leu				
515								520							

&lt;210&gt; 25

&lt;211&gt; 524

&lt;212&gt; PRT

&lt;213&gt; Lyssavirus sp.

&lt;220&gt;

&lt;223&gt; USA7-BT

&lt;400&gt; 25

Met	Ile	Pro	Gln	Ala	Leu	Leu	Phe	Val	Pro	Leu	Leu	Ile	Pro	Ser	Leu
1					5					10				15	

Cys	Leu	Gly	Glu	Phe	Pro	Ile	Tyr	Thr	Ile	Pro	Asp	Lys	Leu	Gly	Pro
20									25					30	

Trp	Thr	Pro	Ile	Asp	Ile	His	His	Leu	Ser	Cys	Pro	Asn	Asn	Leu	Val
35								40				45			

Ala	Glu	Asn	Asp	Gly	Cys	Thr	Ser	Leu	Ser	Gly	Phe	Ser	Tyr	Met	Glu
50								55				60			

Leu	Lys	Val	Gly	Tyr	Ile	Ser	Ala	Ile	Lys	Val	Asn	Gly	Phe	Thr	Cys
65								70				75			80

Thr	Gly	Val	Val	Thr	Glu	Ala	Glu	Thr	Tyr	Thr	Asn	Phe	Val	Gly	Tyr
85									90				95		

Val	Thr	Thr	Thr	Phe	Lys	Arg	Lys	His	Phe	Arg	Pro	Met	Pro	Asp	Ala
100								105				110			

Cys	Arg	Ala	Ala	His	Asp	Trp	Lys	Met	Ala	Gly	Asp	Pro	Arg	Tyr	Glu
115								120				125			

Asp Ser Leu Gln Asn Pro Tyr Pro Asp Tyr His Trp Leu Arg Thr Val  
 130 135 140  
 Lys Thr Thr Lys Glu Ser Leu Val Ile Ile Ser Pro Ser Val Ala Asp  
 145 150 155 160  
 Leu Asp Pro Tyr Asp Lys Ser Leu His Ser Arg Val Phe Pro Ser Gly  
 165 170 175  
 Lys Cys Leu Gly Ile Thr Val Ser Ser Thr Tyr Cys Ser Thr Asn His  
 180 185 190  
 Asp Tyr Thr Ile Trp Met Pro Val Glu Pro Arg Leu Gly Thr Ser Cys  
 195 200 205  
 Asp Ile Phe Thr Asn Ser Arg Gly Lys Arg Ala Ser Lys Gly Gly Arg  
 210 215 220  
 Val Cys Gly Phe Val Asp Glu Arg Gly Leu Tyr Lys Ser Leu Lys Gly  
 225 230 235 240  
 Ala Cys Lys Leu Lys Leu Cys Gly Val Pro Gly Ile Arg Leu Met Asp  
 245 250 255  
 Gly Thr Trp Val Ser Ile Gln Thr Ser Glu Asp Ile Lys Trp Cys Pro  
 260 265 270  
 Pro Asp Arg Leu Val Asn Leu His Asp Phe His Ser Asp Glu Leu Glu  
 275 280 285  
 His Leu Val Val Glu Glu Leu Ile Lys Arg Arg Glu Asn Cys Leu Asp  
 290 295 300  
 Ala Leu Glu Ser Ile Met Thr Thr Lys Ser Val Ser Phe Arg Arg Leu  
 305 310 315 320  
 Ser His Leu Arg Arg Leu Val Pro Gly Phe Gly Lys Ala Tyr Thr Ile  
 325 330 335  
 Phe Asn Lys Thr Leu Ile Glu Ala Asp Ala His Tyr Lys Ser Ile Lys  
 340 345 350  
 Thr Trp Asn Glu Val Ile Pro Ser Lys Gly Cys Leu Glu Val Gly Gly  
 355 360 365  
 Lys Cys His Pro Pro Val Asn Gly Val Phe Phe Asn Gly Ile Ile Leu  
 370 375 380  
 Gly Pro Asp Gly Asn Val Leu Ile Pro Glu Met Gln Ser Ser Leu Leu  
 385 390 395 400  
 Gln Gln His Met Glu Leu Leu Glu Ser Ser Val Ile Pro Leu Met His  
 405 410 415  
 Pro Leu Ala Asp Pro Ser Thr Val Phe Lys Glu Gly Asp Glu Ala Glu  
 420 425 430

Asp Phe Val Glu Val His Leu Pro Asp Val His Lys Arg Ile Ser Gly			
435	440	445	
Val Asp Leu Gly Leu Pro Ser Trp Gly Lys Tyr Leu Leu Met Ser Ala			
450	455	460	
Gly Ala Leu Ala Ile Leu Ile Leu Ala Ile Val Leu Ile Ile Cys Cys			
465	470	475	480
Arg Arg Val Asn Lys Thr Gly Ser Thr Gln Arg Gly His Arg Glu Ser			
485	490	495	
Arg Gly Lys Met Ser Val Ala Pro Gln Asn Gly Lys Ile Ile Ser Ser			
500	505	510	
Trp Glu Leu Tyr Lys Arg Glu Ser Glu Thr Gly Leu			
515	520		

<210> 26  
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 <212> PRT  
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<220>  
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<400> 26  
 Met Leu Leu Gln Ile Val Leu Leu Met Ser Leu Met Val Phe Ser Pro  
 1 5 10 15

Cys Pro Gly Lys Phe Pro Leu Tyr Thr Ile Pro Asp Lys Leu Gly Pro  
 20 25 30

Trp Ser Pro Ile Asp Ile His His Leu Ser Cys Pro Asn Asn Leu Ile  
 35 40 45

Val Glu Asp Glu Gly Cys Thr Ser Leu Ser Gly Phe Ser Tyr Met Glu  
 50 55 60

Leu Lys Val Gly Phe Ile Thr Thr Ile Lys Val Ser Gly Phe Thr Cys  
 65 70 75 80

Thr Gly Val Val Thr Glu Ser Glu Thr Tyr Thr Asn Phe Phe Gly Tyr  
 85 90 95

Val Thr Thr Thr Phe Lys Arg Lys His Phe Arg Pro Thr Pro Glu Phe  
 100 105 110

Cys Arg Asn Ala Tyr Asn Trp Lys Val Ala Gly Asp Pro Arg Tyr Glu  
 115 120 125

Glu Ser Leu His Asn Pro Tyr Pro Asp Tyr His Trp Leu Arg Thr Val  
 130 135 140

Thr Thr Thr Lys Glu Ser Leu Leu Ile Ile Ser Pro Ser Val Val Asp  
 145 150 155 160

Met Asp Pro Tyr Asp Lys Ser Leu His Ser Lys Met Phe Pro Lys Gly  
 165 170 175

Thr Cys Ser Gly Ala Ser Val Pro Ser Ile Phe Cys Ser Thr Asn His  
 180 185 190

Asp Tyr Thr Leu Trp Met Pro Glu Asn Pro Lys Pro Gly Met Ser Cys  
 195 200 205

Asp Ile Phe Thr Thr Ser Lys Gly Lys Lys Ala Ser Lys Gly Gly Lys  
 210 215 220

Val Cys Gly Phe Val Asp Glu Arg Gly Leu Tyr Lys Ser Leu Lys Gly  
 225 230 235 240

Ala Cys Lys Leu Lys Leu Cys Gly Ile Ser Gly Leu Arg Leu Met Asp  
 245 250 255

Gly Ser Trp Val Ser Ile Gln Asn His Glu Glu Ala Lys Trp Cys Ser  
 260 265 270

Pro Asp Gln Leu Val Asn Ile His Asp Phe His Ser Asp Glu Ile Glu  
 275 280 285

His Leu Ile Val Glu Glu Leu Val Arg Lys Arg Glu Glu Cys Leu Asp  
 290 295 300

Ala Leu Glu Ser Ile Met Thr Thr Lys Ser Val Ser Phe Arg Arg Leu  
 305 310 315 320

Ser His Leu Arg Lys Leu Val Pro Gly Phe Gly Lys Ala Tyr Thr Ile  
 325 330 335

Val Asn Lys Thr Leu Met Glu Ala Asp Ala His Tyr Asn Gln Val Arg  
 340 345 350

Thr Trp Asn Glu Ile Ile Pro Ser Lys Gly Cys Leu Lys Val Arg Glu  
 355 360 365

Arg Cys His Pro Pro Tyr Asn Gly Val Phe Phe Asn Gly Ile Ile Leu  
 370 375 380

Ser Pro Asp Gly His Val Leu Ile Pro Glu Met Gln Ser Ser Leu Leu  
 385 390 395 400

Gln Gln His Ile Glu Leu Leu Glu Ser Ser Val Ile Pro Leu Ile His  
 405 410 415

Pro Leu Ala Asp Pro Ser Thr Val Phe Tyr Arg Asp Asp Glu Ala Glu  
 420 425 430

Asp Phe Ile Glu Val His Leu Pro Asp Val Gln Lys Gln Val Ser Gly  
 435 440 445

Ile Asp Leu Gly Leu Ser Glu Trp Glu Arg Tyr Leu Ile Ile Gly Ala  
 450 455 460

Ser Ala Val Ile Leu Phe Ala Leu Ala Ile Ile Phe Ala Val Cys Cys  
 465 470 475 480

Arg Arg Cys Lys Arg Arg Lys Lys Ala Arg Thr Asp Arg Ile Glu Leu  
 485 490 495

Asp Arg Lys Val Ser Val Thr Ser Gln Ser Gly Lys Val Ile Pro Ser  
 500 505 510

Trp Glu Ser Tyr Lys Leu Pro Lys Ser His Phe Arg Ser  
 515 520 525

<210> 27  
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 <212> PRT  
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<220>  
 <223> EBL1POL

<400> 27  
 Met Leu Leu Ser Thr Ala Ile Phe Ala Phe Phe Leu Asn Cys Ala Pro  
 1 5 10 15

Cys Leu Ala Lys Phe Pro Ile Tyr Thr Ile Pro Asp Lys Ile Gly Pro  
 20 25 30

Trp Ser Pro Ile Asp Ile Asn His Leu Ser Cys Pro Asn Asn Leu Ile  
 35 40 45

Val Glu Asp Glu Gly Cys Thr Thr Leu Thr Pro Phe Ser Tyr Met Glu  
 50 55 60

Leu Lys Val Gly Tyr Ile Thr Thr Ile Ile Glu Ser Gly Phe Thr Cys  
 65 70 75 80

Thr Gly Val Ile Thr Glu Ala Glu Thr Tyr Thr Asn Phe Val Gly Tyr  
 85 90 95

Val Thr Thr Phe Lys Arg Lys His Phe Arg Pro Thr Val Ser Ala  
 100 105 110

Cys Arg Asp Ala Tyr Asn Trp Lys Ile Thr Gly Asp Pro Arg Tyr Glu  
 115 120 125

Glu Ser Leu His Asn Pro Tyr Pro Asp Ser His Trp Leu Arg Thr Val  
 130 135 140

Lys Thr Thr Lys Glu Ser Leu Leu Ile Ile Ser Pro Ser Val Val Asp  
 145 150 155 160

Met Asp Ala Tyr Asp Lys Asn Leu Tyr Ser Lys Met Phe Pro Asn Gly  
 165 170 175

Lys Cys Leu Ala Ser Pro Pro Ser Ala Ile Cys Cys Pro Thr Asn His  
 180 185 190

Asp Tyr Thr Ile Trp Ile Pro Glu Asn Pro Lys Pro Gly Leu Ser Cys  
 195 200 205

Asp Ile Phe Thr Thr Ser Lys Gly Lys Lys Ala Thr Lys Asp Gly Arg  
 210 215 220

Leu Cys Gly Phe Val Asp Glu Arg Gly Leu Tyr Lys Ser Leu Lys Gly  
 225 230 235 240

Ala Cys Lys Leu Arg Leu Cys Gly Val Pro Gly Met Arg Leu Met Asp  
 245 250 255

Gly Ser Trp Val Ser Leu Gln Lys Thr Glu Ala Pro Glu Trp Cys Ser  
 260 265 270

Pro Asp Gln Leu Val Asn Val His Asp Phe His Thr Asp Glu Ile Glu  
 275 280 285

His Leu Val Val Glu Glu Leu Val Lys Lys Arg Glu Glu Cys Leu Asp  
 290 295 300

Ala Leu Glu Thr Ile Ile Thr Thr Lys Ser Ile Ser Phe Arg Arg Leu  
 305 310 315 320

Ser His Phe Arg Lys Leu Val Pro Gly Phe Gly Lys Ala Tyr Thr Leu  
 325 330 335

Ile Asn Lys Thr Leu Met Glu Ala Asp Ala His Tyr Lys Ser Val Arg  
 340 345 350

Glu Trp Lys Glu Val Ile Pro Ser Lys Gly Cys Leu Met Ala Gly Gly  
 355 360 365

Arg Cys His Pro His Tyr Ser Gly Ile Phe Phe Asn Gly Ile Ile Leu  
 370 375 380

Ser Pro Gly Gly Asp Val Leu Ile Pro Glu Met Gln Ser Ala Leu Leu  
 385 390 395 400

Gln Gln His Ile Glu Leu Leu Glu Ser Ser Met Ile Pro Leu Arg His  
 405 410 415

Pro Leu Ala Asp Pro Ser Thr Val Phe Arg Lys Asp Asp Glu Ala Glu  
 420 425 430

Asp Phe Val Glu Val His Leu Pro Asp Thr Gln Lys Leu Ile Ser Gly  
 435 440 445

Ile Asp Leu Gly Phe Pro Glu Trp Lys Arg Tyr Phe Leu Ile Gly Ile  
 450 455 460

Ser Val Leu Ala Leu Leu Ala Leu Ala Ile Ile Thr Ala Ala Cys Cys  
 465 470 475 480

Lys Arg Phe Lys Arg Arg Arg Pro Lys Pro Asn Pro Ile Glu Leu  
 485 490 495

Ile Arg Lys Val Ser Val Thr Ser Gln Ser Gly Arg Ala Ile Pro Ser  
 500 505 510

Trp Glu Ser Tyr Lys Val Gly Pro Pro Gly Glu Ser  
 515 520

<210> 28  
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 <212> PRT  
 <213> Lyssavirus sp.

<220>  
 <223> EBL1FRA

<400> 28  
 Met Leu Leu Ser Thr Ala Ile Phe Ala Phe Phe Leu Asn Cys Ala Pro  
 1 5 10 15

Cys Leu Gly Lys Phe Pro Ile Tyr Thr Ile Pro Asp Lys Ile Gly Pro  
 20 25 30

Trp Ser Pro Ile Asp Ile Asn His Leu Ser Cys Pro Asn Asn Leu Ile  
 35 40 45

Val Glu Asp Glu Gly Cys Thr Thr Leu Thr Pro Phe Ser Tyr Met Glu  
 50 55 60

Leu Lys Val Gly Tyr Ile Thr Thr Ile Lys Ile Glu Gly Phe Thr Cys  
 65 70 75 80

Thr Gly Val Ile Thr Glu Ala Glu Thr Tyr Thr Asn Phe Val Gly Tyr  
 85 90 95

Val Thr Thr Thr Phe Lys Arg Lys His Phe Arg Pro Thr Val Ser Ala  
 100 105 110

Cys Arg Asp Ala Tyr Asn Trp Lys Ile Thr Gly Asp Pro Arg Tyr Glu  
 115 120 125

Glu Ser Leu His Asn Pro Tyr Pro Asp Ser His Trp Leu Arg Thr Val  
 130 135 140

Lys Thr Thr Lys Glu Ser Leu Leu Ile Ile Ser Pro Ser Val Val Asp  
 145 150 155 160

Met Asp Ala Tyr Asp Lys Asn Leu Tyr Ser Lys Met Phe Pro Asn Gly  
 165 170 175

Lys Cys Leu Ala Ser Pro Pro Ser Ala Thr Cys Cys Pro Thr Asn His  
 180 185 190

Asp Tyr Thr Ile Trp Ile Pro Glu Asn Pro Lys Pro Gly Leu Ser Cys  
 195 200 205

Asp Ile Phe Thr Thr Ser Lys Gly Lys Lys Ala Thr Lys Asp Gly Lys  
 210 215 220

Leu Cys Gly Phe Val Asp Glu Arg Gly Leu Tyr Lys Ser Leu Lys Gly  
 225 230 235 240

Ala Cys Lys Leu Arg Leu Cys Gly Val Pro Gly Met Arg Leu Met Asp  
 245 250 255

Gly Ser Trp Val Ser Leu Gln Lys Thr Glu Ala Pro Glu Trp Cys Ser  
 260 265 270

Pro Asp Arg Leu Val Asn Ile His Asp Phe His Thr Asp Glu Ile Glu  
 275 280 285

His Leu Val Val Glu Glu Leu Val Lys Lys Arg Glu Glu Cys Leu Asp  
 290 295 300

Ala Leu Glu Thr Ile Ile Thr Thr Lys Ser Ile Ser Phe Arg Arg Leu  
 305 310 315 320

Ser His Phe Arg Lys Leu Val Pro Gly Phe Gly Lys Ala Tyr Thr Leu  
 325 330 335

Ile Asn Lys Thr Leu Met Glu Ala Asp Ala His Tyr Lys Ser Val Arg  
 340 345 350

Glu Trp Thr Glu Val Ile Pro Ser Lys Gly Cys Leu Met Ala Gly Gly  
 355 360 365

Arg Cys His Pro His Tyr Ser Gly Ile Phe Phe Asn Gly Ile Ile Leu  
 370 375 380

Ser Pro Gly Gly Asp Val Leu Ile Pro Glu Met Gln Ser Ala Leu Leu  
 385 390 395 400

Gln Gln His Ile Glu Leu Leu Glu Ser Ser Met Ile Pro Leu Arg His  
 405 410 415

Pro Leu Ala Asp Pro Ser Thr Val Phe Lys Arg Asp Asp Glu Ala Glu  
 420 425 430

Asp Phe Val Glu Val His Leu Pro Asp Thr Gln Lys Leu Ile Ser Gly  
 435 440 445

Ile Asp Leu Gly Phe Pro Glu Trp Lys Arg Tyr Phe Leu Ile Gly Ile  
 450 455 460

Ser Val Leu Ala Leu Leu Ala Leu Ala Ile Ile Thr Ala Ala Cys Cys  
 465 470 475 480

Lys Arg Phe Lys Arg Arg Arg Pro Lys Pro Asn Pro Ile Glu Leu  
 485 490 495

Ile Arg Lys Val Ser Val Thr Ser Gln Ser Gly Arg Ala Ile Pro Ser  
 500 505 510

Trp Glu Ser Tyr Lys Val Gly Thr Thr Ser Glu Ser  
 515 520

<210> 29  
 <211> 524  
 <212> PRT  
 <213> Lyssavirus sp.

<220>  
 <223> EBL2FIN

<400> 29

Met Pro Phe Gln Thr Val Leu Ser Ala Leu Leu Ser Ala Leu Thr Leu  
 1 5 10 15

Cys Ala Gly Lys Phe Pro Ile Tyr Thr Ile Pro Asp Lys Leu Gly Pro  
 20 25 30

Trp Ser Pro Ile Asp Ile His His Leu Ser Cys Pro Thr Asn Met Val  
 35 40 45

Val Glu Asp Glu Gly Cys Thr Thr Leu Thr Val Phe Ser Tyr Met Glu  
 50 55 60

Leu Arg Val Gly Tyr Ile Thr Thr Ile Lys Val Asp Gly Phe Thr Cys  
 65 70 75 80

Thr Gly Val Val Thr Glu Ala Glu Thr Tyr Thr Asn Phe Val Gly Tyr  
 85 90 95

Val Thr Thr Phe Lys Arg Lys His Phe Arg Pro Ser Pro Ser Ala  
 100 105 110

Cys Arg Asp Ala Tyr Ser Trp Lys Asn Ala Gly Asp Pro Arg Tyr Glu  
 115 120 125

Glu Ser Leu His Asn Pro Tyr Pro Asp Ser His Trp Leu Arg Thr Val  
 130 135 140

Thr Thr Thr Lys Glu Ser Leu Leu Ile Ile Ser Pro Ser Val Val Asn  
 145 150 155 160

Met Asp Ala Tyr Asp Lys Thr Leu Tyr Ser Lys Ile Phe Leu Asn Gly  
 165 170 175

Lys Cys Ser Gly Val Ser Gln Val Ser Pro Phe Cys Ser Thr Asn His  
 180 185 190

Asp Tyr Thr Ile Trp Met Pro Glu Asn Pro Asn Pro Gly Val Ser Cys  
 195 200 205

Asp Ile Phe Thr Thr Ser Lys Gly Lys Lys Ala Thr Lys Asp Gly Lys  
 210 215 220

Leu Cys Gly Phe Val Asp Glu Arg Gly Leu Tyr Lys Ser Leu Lys Gly  
 225 230 235 240

Ala Cys Lys Leu Lys Leu Cys Gly Ile Ser Gly Met Arg Leu Met Asp  
 245 250 255

Gly Ser Trp Val Ser Ile Gln Asn His Asp Glu Ala Lys Trp Cys Ser  
 260 265 270  
 Pro Asp Gln Leu Val Asn Ile His Asp Phe His Ser Asp Glu Val Glu  
 275 280 285  
 His Leu Ile Ala Glu Glu Leu Val Lys Lys Arg Glu Glu Cys Leu Asp  
 290 295 300  
 Ala Leu Glu Ser Ile Met Thr Thr Lys Ser Ile Ser Phe Arg Arg Leu  
 305 310 315 320  
 Ser His Leu Arg Lys Leu Val Pro Gly Phe Gly Lys Ala Tyr Thr Ile  
 325 330 335  
 Ile Asn Lys Thr Leu Met Glu Ala Asp Ala His Tyr Lys Ser Ile Arg  
 340 345 350  
 Glu Trp Thr Asp Val Ile Pro Ser Lys Gly Cys Leu Met Ala Gly Gly  
 355 360 365  
 Arg Cys Tyr Pro His His Asn Gly Val Phe Phe Asn Gly Ile Ile Leu  
 370 375 380  
 Ser Pro Asp Gly His Val Leu Ile Pro Glu Met Gln Ser Ala Met Leu  
 385 390 395 400  
 Gln Gln His Ile Glu Leu Leu Glu Ser Ser Val Ile Pro Leu Met His  
 405 410 415  
 Pro Leu Ala Asp Pro Ser Thr Ile Phe Lys Lys Asp Asp Gly Ala Glu  
 420 425 430  
 Asp Phe Val Glu Val His Leu Pro Asp Val Gln Lys Gln Ile Ser Gly  
 435 440 445  
 Ile Asp Leu Gly Leu Pro Glu Trp Lys Arg Tyr Phe Leu Ile Gly Val  
 450 455 460  
 Ser Ala Leu Ala Leu Leu Ala Leu Met Ile Phe Ile Ala Ala Cys Cys  
 465 470 475 480  
 Lys Arg Val Lys His Lys Lys Arg Ala Lys Pro Asn Pro Val Glu Leu  
 485 490 495  
 Ile Arg Lys Val Ser Val Thr Ser Gln Ser Gly Arg Pro Ile Pro Ser  
 500 505 510  
 Trp Glu Ser Tyr Lys Val Glu Thr Gly Gly Gln Ser  
 515 520

<210> 30  
 <211> 524  
 <212> PRT  
 <213> Lyssavirus sp.

<220>  
<223> EBL2HOL

&lt;400&gt; 30

Met	Pro	Phe	Gln	Ala	Val	Leu	Ser	Ala	Leu	Leu	Ser	Ala	Leu	Thr	Leu
1															
					5					10					15

Cys	Val	Gly	Lys	Phe	Pro	Ile	Tyr	Thr	Ile	Pro	Asp	Lys	Leu	Gly	Pro
					20				25						30

Trp	Ser	Pro	Ile	Asp	Ile	His	His	Leu	Ser	Cys	Pro	Thr	Asn	Met	Val
					35			40						45	

Val	Glu	Asp	Glu	Gly	Cys	Thr	Thr	Leu	Thr	Val	Phe	Ser	Tyr	Met	Glu
	50				55						60				

Leu	Lys	Val	Gly	Tyr	Ile	Thr	Thr	Ile	Lys	Val	Asn	Glu	Phe	Thr	Cys
	65				70				75					80	

Thr	Gly	Val	Val	Thr	Glu	Ala	Glu	Thr	Tyr	Thr	Asn	Phe	Val	Gly	Tyr
	85								90					95	

Val	Thr	Thr	Phe	Lys	Arg	Lys	Asp	Phe	Arg	Pro	Ser	Pro	Ser	Ala
	100				105						110			

Cys	Arg	Asp	Ala	Tyr	Ser	Cys	Lys	Thr	Ala	Gly	Asp	Pro	Arg	Tyr	Glu
	115					120					125				

Glu	Ser	Leu	His	Asn	Pro	Tyr	Pro	Asp	Ser	His	Trp	Leu	Thr	Cys	Thr
	130				135						140				

Thr	Thr	Thr	Lys	Glu	Ser	Val	Leu	Ile	Ile	Ser	Pro	Ser	Val	Ala	Asp
	145				150				155					160	

Met	Asp	Ala	Tyr	Asp	Lys	Thr	Leu	Tyr	Ser	Lys	Ile	Phe	Leu	Asn	Gly
	165								170					175	

Lys	Cys	Ser	Gly	Val	Ser	Gln	Val	Ser	Pro	Phe	Cys	Ser	Thr	Asn	His
	180				185								190		

Asp	Tyr	Thr	Ile	Trp	Met	Pro	Glu	Asn	Pro	Asn	Pro	Gly	Val	Ser	Cys
	195					200						205			

Asp	Ile	Phe	Thr	Thr	Ser	Lys	Gly	Lys	Lys	Ala	Thr	Lys	Asp	Gly	Lys
	210				215						220				

Leu	Cys	Gly	Phe	Val	Asp	Glu	Arg	Gly	Leu	Tyr	Lys	Ser	Leu	Lys	Gly
	225				230					235				240	

Ala	Cys	Lys	Leu	Lys	Leu	Cys	Gly	Ile	Ser	Gly	Met	Arg	Leu	Met	Asp
	245					250							255		

Gly	Ser	Trp	Val	Ser	Ile	Gln	Asn	His	Asp	Glu	Ala	Lys	Trp	Cys	Ser
	260					265						270			

Pro	Asp	Gln	Leu	Val	Asn	Ile	His	Asp	Phe	His	Ser	Asp	Glu	Val	Glu
	275					280						285			

His	Leu	Ile	Ala	Glu	Glu	Leu	Val	Lys	Lys	Arg	Glu	Glu	Cys	Leu	Asp
290				295						300					
Ala	Leu	Glu	Ser	Ile	Met	Thr	Thr	Lys	Ser	Ile	Ser	Phe	Arg	Arg	Leu
305					310			315							320
Ser	His	Leu	Arg	Lys	Leu	Val	Pro	Gly	Phe	Gly	Lys	Ala	Tyr	Thr	Val
				325				330							335
Ile	Asn	Lys	Thr	Leu	Met	Glu	Ala	Asp	Ala	His	Tyr	Lys	Ser	Ile	Arg
				340				345							350
Glu	Trp	Thr	Asp	Val	Ile	Pro	Ser	Lys	Gly	Cys	Leu	Met	Ala	Gly	Gly
				355			360				365				
Arg	Cys	Tyr	Pro	His	His	Asn	Gly	Val	Phe	Phe	Asn	Gly	Ile	Ile	Leu
				370			375				380				
Ser	Pro	Asp	Gly	His	Val	Leu	Ile	Pro	Glu	Met	Gln	Ser	Ala	Met	Leu
				385			390			395					400
Gln	Gln	His	Ile	Glu	Leu	Leu	Glu	Ser	Ser	Val	Ile	Pro	Leu	Met	His
				405				410							415
Pro	Leu	Ala	Asp	Pro	Ser	Thr	Ile	Phe	Lys	Lys	Asp	Asp	Gly	Ala	Glu
				420				425							430
Asp	Phe	Val	Glu	Val	His	Leu	Pro	Asp	Val	Gln	Lys	Gln	Ile	Ser	Gly
				435				440							445
Ile	Asp	Leu	Gly	Leu	Pro	Glu	Trp	Lys	Arg	Tyr	Phe	Leu	Ile	Gly	Val
				450			455				460				
Ser	Ala	Leu	Ala	Phe	Leu	Ala	Leu	Met	Ile	Phe	Ile	Ala	Ala	Cys	Cys
				465			470			475					480
Arg	Arg	Val	Lys	Arg	Lys	Arg	Ala	Lys	Pro	Asn	Pro	Val	Glu	Leu	
				485				490							495
Ile	Arg	Lys	Val	Ser	Val	Thr	Ser	Gln	Ser	Gly	Arg	Pro	Ile	Pro	Ser
				500				505							510
Trp	Glu	Ser	Tyr	Lys	Val	Glu	Thr	Gly	Gly	Gln	Ser				
				515				520							

<210> 31  
 <211> 533  
 <212> PRT  
 <213> Lyssavirus sp.

<220>  
 <223> Duv1SAF

<400> 31  
 Met Pro Leu Asn Ala Val Ile Phe Thr Leu Leu Leu Arg Cys Ser Ile  
 1 5 10 15

Cys	Leu	Gly	Lys	Phe	Pro	Phe	Tyr	Thr	Ile	Pro	Asp	Lys	Leu	Gly	Pro
20								25					30		
Trp	Ser	Pro	Ile	Asp	Ile	His	His	Leu	Ser	Cys	Pro	Asn	Asn	Leu	Val
35						40						45			
Val	Glu	Asp	Glu	Gly	Cys	Thr	Thr	Leu	Thr	Pro	Phe	Ser	Tyr	Met	Glu
50						55					60				
Leu	Lys	Val	Gly	Tyr	Ile	Thr	Ser	Ile	Lys	Val	Ser	Gly	Phe	Thr	Cys
65						70			75					80	
Thr	Gly	Val	Val	Thr	Glu	Ala	Glu	Thr	Tyr	Thr	Asn	Phe	Val	Gly	Tyr
									85		90			95	
Val	Thr	Thr	Thr	Phe	Arg	Arg	Arg	His	Phe	Arg	Pro	Ser	Val	Asn	Ser
								100		105			110		
Cys	Arg	Asp	Ala	Tyr	Asn	Trp	Lys	Ile	Ala	Gly	Asp	Pro	Arg	Tyr	Glu
								115		120		125			
Glu	Ser	Leu	His	Asn	Pro	Tyr	Pro	Asp	Ser	His	Trp	Leu	Arg	Thr	Val
								130		135		140			
Lys	Thr	Thr	Lys	Glu	Ser	Leu	Leu	Ile	Ile	Ser	Pro	Ser	Val	Ala	Asp
								145		150		155		160	
Met	Asp	Ala	Tyr	Asp	Lys	Lys	Leu	Tyr	Ser	Lys	Met	Phe	Pro	Asn	Gly
								165		170			175		
Arg	Cys	Ser	Glu	Ile	Ser	Pro	Gly	Ser	Pro	Phe	Cys	Pro	Thr	Asn	His
								180		185			190		
Glu	Tyr	Thr	Ile	Trp	Met	Pro	Glu	Ser	Ser	Asn	Pro	Gly	Ile	Ser	Cys
								195		200		205			
Asp	Ile	Phe	Thr	Arg	Ser	Met	Gly	Lys	Lys	Ala	Thr	Lys	Asp	Gly	Gln
								210		215		220			
Leu	Cys	Gly	Phe	Val	Asp	Glu	Arg	Gly	Leu	Tyr	Lys	Ser	Leu	Lys	Gly
								225		230		235		240	
Ala	Cys	Arg	Leu	Arg	Leu	Cys	Gly	Ile	Ser	Gly	Leu	Arg	Leu	Met	Asp
								245		250			255		
Gly	Ser	Trp	Val	Ser	Leu	Pro	Gln	Val	Asn	Asn	Ser	Glu	Trp	Cys	Ser
								260		265			270		
Pro	Asp	Gln	Leu	Val	Asn	Ile	His	Asp	Phe	His	Ser	Asp	Glu	Ile	Glu
								275		280			285		
His	Leu	Val	Ala	Asp	Glu	Leu	Val	Lys	Lys	Arg	Glu	Asp	Cys	Leu	Asp
								290		295		300			
Ala	Leu	Glu	Thr	Ile	Ile	Phe	Thr	Lys	Ser	Ile	Ser	Phe	Arg	Arg	Leu
								305		310		315		320	

Ser Arg Leu Arg Lys Leu Val Pro Gly Phe Gly Lys Ala Tyr Thr Ile  
 325 330 335  
 Ile Asn Arg Thr Leu Met Glu Ala Glu Ala His Tyr Lys Ser Val Arg  
 340 345 350  
 Glu Trp Lys Glu Ile Ile Pro Ser Lys Gly Cys Leu Lys Ala Gly Gly  
 355 360 365  
 Arg Cys Tyr Pro His His Asn Ile Val Phe Phe Asn Gly Ile Ile Leu  
 370 375 380  
 Gly Pro Gly Gly Lys Ile Leu Ile Pro Glu Met Gln Ser Ala Leu Leu  
 385 390 395 400  
 Gln Gln His Ile Glu Leu Leu Glu Ser Ser Val Val Pro Leu Lys His  
 405 410 415  
 Pro Leu Ala Asp Pro Ser Thr Val Phe Lys Asn Asp Asp Glu Ala Glu  
 420 425 430  
 Ser Phe Val Asp Val His Leu Pro Asp Thr Asn Gln Lys Ile Ser Gly  
 435 440 445  
 Ile Asp Leu Gly Leu Pro Glu Trp Lys Arg Tyr Phe Leu Ile Gly Val  
 450 455 460  
 Ser Ala Val Ala Leu Leu Ala Leu Ser Ile Ile Met Arg Val Cys Cys  
 465 470 475 480  
 Lys Arg Phe Lys Asn Arg Arg Lys Ser Lys Pro Ser Pro Val Glu Leu  
 485 490 495  
 Thr Arg Lys Val Ser Val Ile Ser Lys Gly Asn Gly Pro Val Pro Ser  
 500 505 510  
 Trp Glu Ser Tyr Lys Glu Gly Thr Thr Gly Asp Val Arg Asn Thr Thr  
 515 520 525  
 Pro Ser Thr Arg Glu  
 530

<210> 32  
 <211> 533  
 <212> PRT  
 <213> Lyssavirus sp.

<220>  
 <223> Duv2SAF

<400> 32  
 Met Pro Leu Asn Ala Val Ile Phe Thr Leu Leu Leu Arg Cys Ser Ile  
 1 5 10 15  
 Cys Leu Gly Lys Phe Pro Phe Tyr Thr Ile Pro Asp Lys Leu Gly Pro  
 20 25 30

Trp Ser Pro Ile Asp Ile His His Leu Ser Cys Pro Asn Asn Leu Val  
 35 40 45

Val Glu Asp Glu Gly Cys Thr Thr Leu Thr Pro Phe Ser Tyr Met Glu  
 50 55 60

Leu Lys Val Gly Tyr Ile Thr Ser Ile Lys Val Ser Gly Phe Thr Cys  
 65 70 75 80

Thr Gly Val Val Thr Glu Ala Glu Thr Tyr Thr Asn Phe Val Gly Tyr  
 85 90 95

Val Thr Thr Thr Phe Arg Arg Arg His Phe Arg Pro Ser Val Asn Ser  
 100 105 110

Cys Arg Asp Ala Tyr Asn Trp Lys Ile Ala Gly Asp Pro Arg Tyr Glu  
 115 120 125

Glu Ser Leu His Asn Pro Tyr Pro Asp Ser His Trp Leu Arg Thr Val  
 130 135 140

Lys Thr Thr Lys Glu Ser Leu Leu Ile Ile Ser Pro Ser Val Ala Asp  
 145 150 155 160

Met Asp Ala Tyr Asp Lys Lys Leu Tyr Ser Lys Met Phe Pro Asn Gly  
 165 170 175

Arg Cys Ser Glu Ile Ser Pro Gly Ser Pro Phe Cys Pro Thr Asn His  
 180 185 190

Glu Tyr Thr Ile Trp Met Pro Glu Ser Ser Asn Pro Gly Ile Ser Cys  
 195 200 205

Asp Ile Phe Thr Arg Ser Met Gly Lys Lys Ala Thr Lys Asp Gly Gln  
 210 215 220

Leu Cys Gly Phe Val Asp Glu Arg Gly Leu Tyr Lys Ser Leu Lys Gly  
 225 230 235 240

Ala Cys Arg Leu Arg Leu Cys Gly Ile Ser Gly Leu Arg Leu Met Asp  
 245 250 255

Gly Ser Trp Val Ser Leu Pro Gln Val Asn Asn Ser Glu Trp Cys Ser  
 260 265 270

Pro Asp Gln Leu Val Asn Ile His Asp Phe His Ser Asp Glu Ile Glu  
 275 280 285

His Leu Val Ala Asp Glu Leu Val Lys Lys Arg Glu Asp Cys Leu Asp  
 290 295 300

Ala Leu Glu Thr Ile Leu Phe Thr Lys Ser Ile Ser Phe Arg Arg Leu  
 305 310 315 320

Ser His Leu Arg Lys Leu Val Pro Gly Phe Gly Lys Ala Tyr Thr Ile  
 325 330 335

Ile Asn Arg Thr Leu Met Glu Ala Glu Ala His Tyr Lys Ser Val Arg  
 340 345 350  
 Glu Trp Lys Glu Ile Ile Pro Ser Lys Gly Cys Leu Lys Ala Gly Gly  
 355 360 365  
 Arg Cys Tyr Pro His His Asn Gly Ile Phe Phe Asn Gly Ile Ile Leu  
 370 375 380  
 Gly Pro Gly Gly Glu Ile Leu Ile Pro Glu Met Gln Ser Ala Leu Leu  
 385 390 395 400  
 Gln Gln His Ile Glu Leu Leu Glu Ser Ser Val Val Pro Leu Lys His  
 405 410 415  
 Pro Leu Ala Asp Pro Ser Thr Val Phe Lys Asn Asp Asp Glu Ala Glu  
 420 425 430  
 Ser Phe Val Asp Val His Leu Pro Asp Thr Asn Gln Lys Ile Ser Gly  
 435 440 445  
 Ile Asp Leu Gly Leu Pro Glu Trp Lys Arg Tyr Phe Leu Ile Gly Val  
 450 455 460  
 Ser Ala Val Ala Leu Leu Ala Leu Ser Ile Ile Ala Val Cys Cys  
 465 470 475 480  
 Lys Arg Phe Arg Lys Arg Lys Lys Ser Lys Pro Gly Pro Val Glu Leu  
 485 490 495  
 Thr Arg Lys Val Ser Val Ile Ser Lys Gly Asn Gly Pro Val Pro Ser  
 500 505 510  
 Trp Glu Ser Tyr Lys Glu Gly Thr Thr Gly Asp Val Arg Asn Thr Thr  
 515 520 525  
 Pro Ser Thr Arg Glu  
 530

<210> 33  
 <211> 522  
 <212> PRT  
 <213> Lyssavirus sp.

<220>  
 <223> Lag1NGA

<400> 33  
 Met Ser Gln Leu Asn Leu Ile Leu Phe Phe Cys Val Ile Ile Val Leu  
 1 5 10 15  
 Ser Val Glu Asp Phe Pro Leu Tyr Thr Ile Pro Glu Lys Ile Gly Pro  
 20 25 30  
 Trp Thr Pro Ile Asp Leu Ile His Leu Ser Cys Pro Asn Asn Leu Gln  
 35 40 45

Ser Glu Asp Glu Gly Cys Gly Thr Ser Ser Ser Val Ser Tyr Val Glu  
 50 55 60

Leu Lys Thr Gly Tyr Leu Thr His Gln Lys Val Ser Gly Phe Thr Cys  
 65 70 75 80

Thr Gly Val Val Asn Glu Ala Val Thr Tyr Thr Asn Phe Val Gly Tyr  
 85 90 95

Val Thr Thr Thr Phe Lys Arg Lys His Phe Lys Pro Thr Ala Leu Ala  
 100 105 110

Cys Arg Asp Ala Tyr His Trp Lys Ile Ser Gly Asp Pro Arg Tyr Glu  
 115 120 125

Glu Ser Leu His Thr Pro Tyr Pro Asp Asn Ser Trp Leu Arg Thr Val  
 130 135 140

Thr Thr Thr Lys Glu Ser Leu Val Ile Ile Ser Pro Ser Ile Val Glu  
 145 150 155 160

Met Asp Val Tyr Ser Arg Thr Leu His Ser Pro Met Phe Pro Thr Gly  
 165 170 175

Thr Cys Ser Arg Phe Tyr Pro Ser Ser Pro Ser Cys Ala Thr Asn His  
 180 185 190

Asp Tyr Thr Leu Trp Leu Pro Asp Asp Pro Asn Leu Ser Leu Ala Cys  
 195 200 205

Asp Ile Phe Val Thr Ser Thr Gly Lys Lys Ser Met Asn Gly Ser Arg  
 210 215 220

Met Cys Gly Phe Thr Asp Glu Arg Gly Tyr Tyr Arg Thr Ile Lys Gly  
 225 230 235 240

Ala Cys Lys Leu Thr Leu Cys Gly Lys Pro Gly Leu Arg Leu Phe Asp  
 245 250 255

Gly Thr Trp Ile Ser Phe Thr Arg Pro Glu Val Thr Thr Trp Cys Leu  
 260 265 270

Pro Asn Gln Leu Val Asn Ile His Asn Asn Arg Ile Asp Glu Val Glu  
 275 280 285

His Leu Ile Val Glu Asp Leu Ile Arg Lys Arg Glu Glu Cys Leu Asp  
 290 295 300

Thr Leu Glu Thr Val Leu Met Ser Lys Ser Ile Ser Phe Arg Arg Val  
 305 310 315 320

Ser His Phe Arg Lys Leu Val Pro Gly Tyr Gly Lys Ala Tyr Thr Ile  
 325 330 335

Leu Asn Gly Ser Leu Ile Gln Thr Asn Val His Tyr Leu Lys Val Asp  
 340 345 350

Asn Trp Ser Glu Ile Leu Pro Ser Lys Gly Cys Leu Lys Ile Asn Asn  
 355 360 365  
 Gln Cys Val Ala His Asp Glu Gly Val Phe Phe Asn Gly Ile Ile Lys  
 370 375 380  
 Gly Pro Asp Gly His Ile Leu Ile Pro Glu Met Gln Ser Ser Leu Trp  
 385 390 395 400  
 Lys Gln His Met Asp Leu Phe Lys Ala Ala Val Phe Pro Leu Arg His  
 405 410 415  
 Pro Leu Ile Glu Pro Gly Ser Leu Phe Asn Lys Asp Gly Asp Ala Asp  
 420 425 430  
 Glu Phe Val Asp Val His Met Pro Asp Val His Lys Leu Val Ser Asp  
 435 440 445  
 Val Asp Leu Gly Leu Pro Asp Trp Ser Leu Tyr Ala Leu Ile Gly Ala  
 450 455 460  
 Thr Ile Ile Ala Phe Phe Ile Leu Ile Cys Leu Ile Arg Ile Cys Cys  
 465 470 475 480  
 Lys Lys Arg Gly Arg Arg Asn Ser Pro Thr Asn Arg Pro Asp Leu Pro  
 485 490 495  
 Ile Gly Leu Ser Thr Thr Pro Gln Pro Lys Ser Lys Val Ile Ser Ser  
 500 505 510  
 Trp Glu Ser Tyr Lys Gly Thr Ser Asn Val  
 515 520

<210> 34  
 <211> 522  
 <212> PRT  
 <213> Lyssavirus sp.

<220>  
 <223> Lag2CAR

<400> 34  
 Met Ser Gln Leu Ile Leu Ile Pro Phe Leu Cys Val Val Ile Val Ile  
 1 5 10 15

Ser Val Gly Asp Phe Pro Leu Tyr Thr Ile Pro Glu Lys Ile Gly Thr  
 20 25 30

Trp Thr Pro Ile Asp Leu Ile His Leu Ser Cys Pro Asn Asn Leu Leu  
 35 40 45

Ser Glu Asp Asp Gly Cys Ser Asn Thr Ala Thr Phe Asn Tyr Ile Glu  
 50 55 60

Leu Lys Thr Gly Tyr Leu Thr His Gln Lys Val Ser Gly Phe Thr Cys  
 65 70 75 80

Thr Gly Val Val Asn Glu Ala Val Thr Tyr Thr Asn Phe Val Gly Tyr  
 85 90 95  
 Val Thr Thr Thr Phe Lys Arg Lys His Phe Lys Pro Thr Ala Leu Ala  
 100 105 110  
 Cys Arg Asp Ala Phe His Trp Lys Ile Ser Gly Asp Pro Arg Tyr Glu  
 115 120 125  
 Glu Ser Leu His Thr Pro Tyr Pro Asp Asn Ser Trp Leu Arg Thr Val  
 130 135 140  
 Thr Thr Thr Lys Glu Ser Leu Leu Ile Ile Ser Pro Ser Ile Val Glu  
 145 150 155 160  
 Met Asp Val Tyr Ser Arg Thr Leu His Ser Pro Met Phe Pro Gly Gly  
 165 170 175  
 Val Cys Ser Lys Phe Tyr Pro Ser Ser Pro Ser Cys Pro Thr Asn His  
 180 185 190  
 Asp Tyr Thr Leu Trp Leu Pro Glu Asp Ala Asn Leu Ser Met Ala Cys  
 195 200 205  
 Asp Ile Phe Ile Thr Ser Thr Gly Lys Lys Ser Met Asn Gly Ser Arg  
 210 215 220  
 Met Cys Gly Phe Thr Asp Glu Arg Gly Phe Tyr Arg Thr Leu Lys Gly  
 225 230 235 240  
 Ala Cys Lys Leu Thr Leu Cys Gly Lys Pro Gly Leu Arg Leu Tyr Asp  
 245 250 255  
 Gly Thr Trp Val Ser Phe Thr Arg Pro Glu Ile Asn Val Trp Cys Ser  
 260 265 270  
 Pro Asn Gln Leu Val Asn Val His Asn Asn Arg Leu Asp Glu Ile Glu  
 275 280 285  
 His Leu Ile Val Gly Asp Leu Ile Arg Lys Arg Glu Glu Cys Leu Asp  
 290 295 300  
 Thr Leu Glu Thr Ile Leu Met Ser Lys Ser Ile Ser Phe Arg Arg Leu  
 305 310 315 320  
 Ser His Phe Arg Lys Leu Val Pro Gly Tyr Gly Lys Ala Tyr Thr Ile  
 325 330 335  
 Ile Asn Gly Ser Leu Met Glu Thr Asn Val His Tyr Leu Arg Val Asp  
 340 345 350  
 Ser Trp Asn Asp Ile Leu Pro Ser Lys Gly Cys Leu Lys Met Asn Lys  
 355 360 365  
 Gln Cys Val Asp Ser Tyr Arg Gly Val Phe Phe Asn Gly Ile Ile Lys  
 370 375 380

Gly His Asp Gly His Ile Leu Ile Pro Glu Met Gln Ser Ser Leu Leu  
 385 390 395 400  
 Lys Gln His Met Asp Leu Leu Lys Ala Ala Val Phe Pro Leu Arg His  
 405 410 415  
 Pro Leu Ile Asp Gln Asn Ser Leu Phe Lys Lys Asp Gly Asp Ala Asp  
 420 425 430  
 Asp Phe Val Glu Val His Met Pro Asp Ile Gln Lys Leu Ile Ser Asp  
 435 440 445  
 Val Asp Leu Gly Leu Pro Ser Trp Gly Leu Tyr Val Met Ile Gly Ala  
 450 455 460  
 Ala Val Ile Ala Phe Leu Val Leu Ile Cys Leu Ile Arg Ile Cys Cys  
 465 470 475 480  
 Lys Lys Lys Thr Arg Thr Arg Thr Ser Met Glu Arg Pro Asp Pro Pro  
 485 490 495  
 Ile Ser Leu Ser Thr Thr Pro Gln Ser Arg Ala Lys Val Val Ser Ser  
 500 505 510  
 Trp Glu Ser Tyr Lys Gly Ser Ser Asn Val  
 515 520

<210> 35  
 <211> 523  
 <212> PRT  
 <213> Lyssavirus sp.

<220>  
 <223> Mok3ETP

<400> 35  
 Met Asn Ile Pro Cys Phe Ala Val Ile Leu Ser Leu Ala Thr Thr His  
 1 5 10 15  
 Cys Leu Glu Lys Phe Leu Ile Tyr Thr Ile Pro Glu Lys Ile Glu Lys  
 20 25 30  
 Trp Thr Pro Ile Asp Met Ile His Leu Ser Cys Pro Asn Asn Met Leu  
 35 40 45  
 Ser Glu Glu Glu Gly Cys Asn Thr Glu Ser Pro Phe Thr Tyr Phe Glu  
 50 55 60  
 Leu Lys Ser Gly Tyr Leu Ala His Gln Lys Val Pro Gly Phe Thr Cys  
 65 70 75 80  
 Thr Gly Val Val Asn Glu Ala Glu Thr Tyr Thr Asn Phe Val Gly Tyr  
 85 90 95  
 Val Thr Thr Thr Phe Lys Arg Lys His Phe Lys Pro Thr Val Ala Ala  
 100 105 110

Cys Arg Asp Ala Tyr Asn Trp Lys Val Ser Gly Asp Pro Arg Tyr Glu  
 115 120 125

Glu Ser Leu His Thr Pro Tyr Pro Asp Ser Ser Trp Leu Arg Thr Val  
 130 135 140

Thr Thr Thr Lys Glu Ala Leu Leu Ile Ile Ser Pro Ile Val Glu Asp  
 145 150 155 160

Met Ile Ala Gly Arg Lys Thr Leu His Ser Pro Met Phe Pro Ser Gly  
 165 170 175

Lys Cys Ser Lys Leu Tyr Pro Ser Val Pro Ser Cys Thr Thr Asn His  
 180 185 190

Asp Tyr Thr Leu Trp Leu Pro Glu Asp Ser Ser Leu Ser Leu Ile Cys  
 195 200 205

Asp Ile Phe Thr Ser Ser Gly Gln Lys Ala Met Asn Gly Ser Arg  
 210 215 220

Ile Cys Gly Phe Lys Asp Glu Arg Gly Phe Tyr Arg Ser Leu Lys Gly  
 225 230 235 240

Ser Cys Lys Leu Thr Leu Cys Gly Lys Pro Gly Ile Arg Leu Phe Asp  
 245 250 255

Gly Thr Trp Val Ser Phe Thr Lys Pro Asp Val His Val Trp Cys Thr  
 260 265 270

Pro Asn Gln Leu Val Asn Ile His Asn Asp Arg Leu Asp Glu Val Glu  
 275 280 285

His Leu Ile Val Asp Asp Ile Ile Lys Lys Arg Glu Glu Cys Leu Asp  
 290 295 300

Thr Leu Glu Thr Ile Leu Met Ser Gln Ser Val Ser Phe Arg Arg Leu  
 305 310 315 320

Ser His Phe Arg Lys Leu Val Pro Gly Tyr Gly Lys Ala Tyr Thr Ile  
 325 330 335

Leu Asn Gly Ser Leu Met Glu Thr Asn Val Tyr Tyr Lys Arg Val Asp  
 340 345 350

Arg Trp Ala Asp Ile Leu Pro Ser Arg Gly Cys Leu Lys Val Gly Gln  
 355 360 365

Gln Cys Met Asp Pro Val Lys Asn Leu Val Phe Phe Asn Gly Ile Ile  
 370 375 380

Lys Gly Pro Asp Gly Gln Ile Leu Ile Pro Glu Met Gln Ser Glu Gln  
 385 390 395 400

Leu Lys Gln His Met Asp Leu Leu Lys Ala Ala Met Phe Pro Leu Arg  
 405 410 415

His Pro Leu Ile Asn Arg Glu Ala Val Phe Lys Lys Asp Gly Asn Ala  
 420 425 430

Asp Asp Phe Val Asp Leu His Met Pro Asp Val Gln Lys Ser Val Ser  
 435 440 445

Asp Val Asp Leu Gly Leu Pro His Trp Gly Phe Trp Leu Leu Val Gly  
 450 455 460

Ala Thr Val Val Ala Phe Val Val Leu Ala Cys Leu Leu Arg Val Cys  
 465 470 475 480

Cys Arg Arg Met Arg Arg Arg Ser Leu Arg Ala Thr Gln Asp Ile  
 485 490 495

Pro Leu Ser Val Ala Pro Ala Pro Val Pro Arg Ala Lys Val Val Ser  
 500 505 510

Ser Trp Glu Ser Ser Lys Gly Leu Pro Gly Thr  
 515 520

<210> 36  
 <211> 523  
 <212> PRT  
 <213> Lyssavirus sp.

<220>  
 <223> Mok2ZIM

<400> 36  
 Met Asn Ile Pro Cys Phe Val Val Ile Leu Ser Leu Ala Thr Thr His  
 1 5 10 15

Ser Leu Gly Glu Phe Pro Leu Tyr Thr Ile Pro Glu Lys Ile Glu Lys  
 20 25 30

Trp Thr Pro Ile Asp Met Ile His Leu Ser Cys Pro Asn Asn Leu Leu  
 35 40 45

Ser Glu Glu Glu Gly Cys Asn Ala Glu Ser Ser Phe Thr Tyr Phe Glu  
 50 55 60

Leu Lys Ser Gly Tyr Leu Ala His Gln Lys Val Pro Gly Phe Thr Cys  
 65 70 75 80

Thr Gly Val Val Asn Glu Ala Glu Thr Tyr Thr Asn Phe Val Gly Tyr  
 85 90 95

Val Thr Thr Phe Lys Arg Lys His Phe Arg Pro Thr Val Ala Ala  
 100 105 110

Cys Arg Asp Ala Tyr Asn Trp Lys Val Ser Gly Asp Pro Arg Tyr Glu  
 115 120 125

Glu Ser Leu His Thr Pro Tyr Pro Asp Ser Ser Trp Leu Arg Thr Val  
 130 135 140

Thr Thr Thr Lys Glu Ser Leu Leu Ile Ile Ser Pro Ser Ile Val Glu  
 145 150 155 160

Met Asp Ile Tyr Gly Arg Thr Leu His Ser Pro Met Phe Pro Ser Gly  
 165 170 175

Val Cys Ser Asn Val Tyr Pro Ser Val Pro Ser Cys Glu Thr Asn His  
 180 185 190

Asp Tyr Thr Leu Trp Leu Pro Glu Asp Pro Ser Leu Ser Leu Val Cys  
 195 200 205

Asp Ile Phe Thr Ser Ser Asn Gly Lys Lys Ala Met Asn Gly Ser Arg  
 210 215 220

Ile Cys Gly Phe Lys Asp Glu Arg Gly Phe Tyr Arg Ser Leu Lys Gly  
 225 230 235 240

Ala Cys Lys Leu Thr Leu Cys Gly Arg Pro Gly Ile Arg Leu Phe Asp  
 245 250 255

Gly Thr Trp Val Ser Phe Thr Lys Pro Asp Val His Val Lys Thr Cys  
 260 265 270

Asn Pro Asp Ile Leu Val Asn Ile His Asn Asp Arg Leu Asp Glu Ile  
 275 280 285

Glu His Leu Ile Val Glu Asp Ile Ile Lys Lys Arg Glu Glu Cys Leu  
 290 295 300

Asp Thr Leu Glu Thr Ile Leu Met Ser Gln Ser Val Ser Phe Arg Arg  
 305 310 315 320

Leu Ser His Phe Arg Lys Leu Val Pro Gly Tyr Gly Lys Ala Tyr Thr  
 325 330 335

Ile Leu Asn Gly Ser Leu Met Glu Thr Asn Val Tyr Tyr Lys Arg Val  
 340 345 350

Asp Lys Trp Ala Asp Ile Leu Pro Ser Lys Gly Cys Leu Lys Val Gly  
 355 360 365

Gln Gln Cys Met Glu Pro Val Lys Gly Val Leu Phe Asn Gly Ile Ile  
 370 375 380

Lys Gly Pro Asp Gly Gln Ile Leu Ile Pro Glu Met Gln Ser Glu Gln  
 385 390 395 400

Leu Lys Gln His Met Asp Leu Leu Lys Ala Ala Val Phe Pro Leu Arg  
 405 410 415

His Pro Leu Ile Ser Arg Glu Ala Val Phe Lys Lys Asp Gly Asp Ala  
 420 425 430

Asp Asp Phe Val Asp Leu His Met Pro Asp Val His Lys Ser Val Ser  
 435 440 445

Asp Val Asp Leu Gly Leu Pro His Trp Gly Phe Trp Met Leu Ile Gly  
 450 455 460

Ala Thr Ile Val Ala Phe Val Val Leu Val Cys Leu Leu Arg Val Cys  
 465 470 475 480

Cys Lys Arg Val Arg Arg Arg Ser Gly Arg Ala Thr Gln Glu Ile  
 485 490 495

Pro Leu Ser Phe Pro Ser Ala Pro Val Pro Arg Ala Lys Val Val Ser  
 500 505 510

Ser Trp Glu Ser Tyr Lys Gly Leu Pro Gly Thr  
 515 520

<210> 37

<211> 129

<212> PRT

<213> Lyssavirus sp.

<220>

<223> Lag3SAF

<400> 37

Cys Leu Asp Thr Leu Glu Thr Ile Leu Met Ser Lys Ser Ile Ser Phe  
 1 5 10 15

Arg Arg Leu Ser His Phe Arg Lys Leu Val Pro Gly Tyr Gly Lys Ala  
 20 25 30

Tyr Thr Ile Ile Asn Gly Ser Leu Met Glu Thr Asn Val His Tyr Leu  
 35 40 45

Arg Val Asp Ser Trp Asn Asp Ile Leu Pro Ser Lys Gly Cys Leu Lys  
 50 55 60

Met Asn Lys Gln Cys Val Asp Ser Tyr Arg Gly Val Phe Phe Asn Gly  
 65 70 75 80

Ile Ile Lys Gly Leu Asp Gly His Ile Leu Ile Pro Glu Met Gln Ser  
 85 90 95

Ser Leu Leu Lys Gln His Met Asp Leu Leu Lys Ala Ala Val Phe Pro  
 100 105 110

Leu Arg His Pro Leu Ile Asp Gln Asn Ser Leu Phe Lys Lys Asp Gly  
 115 120 125

Asp

<210> 38

<211> 129

<212> PRT

<213> Lyssavirus sp.

&lt;220&gt;

&lt;223&gt; Lag4SAF

&lt;400&gt; 38

Cys	Leu	Asp	Thr	Leu	Glu	Thr	Ile	Leu	Met	Ser	Lys	Ser	Ile	Ser	Phe
1				5				10							15

Arg	Arg	Leu	Ser	His	Phe	Arg	Lys	Leu	Val	Pro	Gly	Tyr	Gly	Lys	Ala
					20			25						30	

Tyr	Thr	Ile	Ile	Asn	Gly	Ser	Leu	Met	Glu	Thr	Asn	Val	His	Tyr	Leu
					35			40					45		

Arg	Val	Asp	Ser	Trp	Asn	Asp	Ile	Leu	Pro	Ser	Lys	Gly	Cys	Leu	Lys
						50		55			60				

Met	Asn	Lys	Gln	Cys	Val	Asp	Ser	Tyr	Arg	Gly	Val	Phe	Phe	Asn	Gly
					65		70		75				80		

Ile	Ile	Lys	Gly	Leu	Asp	Gly	His	Ile	Leu	Ile	Pro	Glu	Met	Gln	Ser
					85				90				95		

Ser	Leu	Leu	Lys	Gln	His	Met	Asp	Leu	Leu	Lys	Ala	Ala	Val	Phe	Pro
						100		105					110		

Leu	Arg	His	Pro	Ile	Asp	Gln	Asn	Ser	Leu	Phe	Lys	Lys	Asp	Gly
						115		120			125			

Asp

&lt;210&gt; 39

&lt;211&gt; 129

&lt;212&gt; PRT

&lt;213&gt; Lyssavirus sp.

&lt;220&gt;

&lt;223&gt; Mok1SAF

&lt;400&gt; 39

Cys	Leu	Asp	Thr	Leu	Glu	Thr	Val	Phe	Met	Ser	Gln	Ser	Val	Ser	Phe
1				5				10						15	

Arg	Arg	Leu	Ser	His	Phe	Arg	Lys	Leu	Val	Pro	Gly	Tyr	Gly	Lys	Ala
					20			25					30		

Tyr	Thr	Ile	Leu	Asn	Gly	Ser	Leu	Met	Glu	Ala	Asn	Val	Tyr	Tyr	Lys
					35			40					45		

Arg	Val	Asp	Arg	Trp	Ala	Asp	Ile	Leu	Pro	Ser	Lys	Gly	Cys	Leu	Lys
						50		55			60				

Val	Gly	Gln	Gln	Cys	Met	Asp	Pro	Val	Asn	Gly	Val	Leu	Phe	Asn	Gly
					65		70		75				80		

Ile	Ile	Lys	Gly	Pro	Asp	Gly	Gln	Ile	Leu	Ile	Pro	Glu	Met	Gln	Ser
					85				90				95		

Glu Gln Leu Lys Gln His Met Asp Leu Leu Lys Ala Ala Val Phe Pro  
100 105 110

Leu Arg His Pro Leu Ile Ser Gln Glu Ala Val Phe Lys Lys Asp Gly  
115 120 125

Asp

<210> 40

<211> 522

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Consensus  
sequence

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<223> variable amino acid

<400> 40

Met Xaa Leu Gln Ala Val Ile Phe Xaa Leu Leu Leu Xaa Xaa Xaa Leu  
1 5 10 15

Cys Leu Gly Lys Phe Pro Ile Tyr Thr Ile Pro Asp Lys Leu Gly Pro  
20 25 30

Trp Ser Pro Ile Asp Ile His His Leu Ser Cys Pro Asn Asn Leu Val  
 35 40 45

Val Glu Asp Glu Gly Cys Thr Thr Leu Xaa Pro Phe Ser Tyr Met Glu  
 50 55 60

Leu Lys Val Gly Tyr Ile Thr Thr Ile Lys Val Ser Gly Phe Thr Cys  
 65 70 75 80

Thr Gly Val Val Thr Glu Ala Glu Thr Tyr Thr Asn Phe Val Gly Tyr  
 85 90 95

Val Thr Thr Thr Phe Lys Arg Lys His Phe Arg Pro Thr Val Xaa Ala  
 100 105 110

Cys Arg Asp Ala Tyr Asn Trp Lys Ile Ala Gly Asp Pro Arg Tyr Glu  
 115 120 125

Glu Ser Leu His Asn Pro Tyr Pro Asp Ser His Trp Leu Arg Thr Val  
 130 135 140

Thr Thr Thr Lys Glu Ser Leu Leu Ile Ile Ser Pro Ser Val Val Asp  
 145 150 155 160

Met Asp Ala Tyr Asp Lys Thr Leu His Ser Lys Met Phe Pro Asn Gly  
 165 170 175

Lys Cys Ser Gly Xaa Ser Pro Ser Pro Phe Cys Xaa Thr Asn His  
 180 185 190

Asp Tyr Thr Ile Trp Met Pro Glu Asn Pro Asn Pro Gly Leu Ser Cys  
 195 200 205

Asp Ile Phe Thr Thr Ser Lys Gly Lys Lys Ala Thr Lys Gly Gly Arg  
 210 215 220

Leu Cys Gly Phe Val Asp Glu Arg Gly Leu Tyr Lys Ser Leu Lys Gly  
 225 230 235 240

Ala Cys Lys Leu Lys Leu Cys Gly Ile Pro Gly Leu Arg Leu Met Asp  
 245 250 255

Gly Ser Trp Val Ser Xaa Gln Xaa Xaa Glu Xaa Xaa Lys Trp Cys Ser  
 260 265 270

Pro Asp Gln Leu Val Asn Ile His Asp Phe His Ser Asp Glu Ile Glu  
 275 280 285

His Leu Ile Val Glu Glu Leu Val Lys Lys Arg Glu Glu Cys Leu Asp  
 290 295 300

Ala Leu Glu Thr Ile Xaa Thr Thr Lys Ser Ile Ser Phe Arg Arg Leu  
 305 310 315 320

Ser His Leu Arg Lys Leu Val Pro Gly Phe Gly Lys Ala Tyr Thr Ile  
 325 330 335

Ile Asn Lys Thr Leu Met Glu Ala Asp Ala His Tyr Lys Ser Val Arg  
 340 345 350

Glu Trp Xaa Glu Ile Ile Pro Ser Lys Gly Cys Leu Lys Ala Gly Gly  
 355 360 365

Arg Cys His Pro His Xaa Asn Gly Val Phe Phe Asn Gly Ile Ile Leu  
 370 375 380

Gly Pro Asp Gly His Val Leu Ile Pro Glu Met Gln Ser Ala Leu Leu  
 385 390 395 400

Gln Gln His Ile Glu Leu Leu Glu Ser Ser Val Ile Pro Leu Arg His  
 405 410 415

Pro Leu Ala Asp Pro Ser Thr Val Phe Lys Lys Asp Asp Glu Ala Glu  
 420 425 430

Asp Phe Val Glu Val His Leu Pro Asp Val Gln Lys Xaa Ile Ser Gly  
 435 440 445

Ile Asp Leu Gly Leu Pro Glu Trp Lys Arg Tyr Phe Leu Ile Gly Ala  
 450 455 460

Ser Ala Leu Ala Leu Leu Ala Leu Ala Ile Ile Ala Val Cys Cys  
 465 470 475 480

Lys Arg Val Lys Arg Arg Arg Xaa Xaa Lys Pro Asn Pro Xaa Glu Leu  
 485 490 495

Ile Arg Lys Val Ser Val Thr Ser Gln Ser Gly Lys Val Ile Pro Ser  
 500 505 510

Trp Glu Ser Tyr Lys Xaa Gly Thr Xaa Gly  
 515 520

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